SEQUENCE LISTING





<110> Innogenetics N.V.

<120> New sequences of hepatitis C virus genotypes for diagnosis, prophylaxis and therapy

<130> 35

<160> 284

<170> PatentIn version 3.1

<210> 1

<211> 213

<212> DNA

<213> hepatitis C virus

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120

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213

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PRT <212>

<213> hepatitis C virus

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Ala Gln Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Pro Thr 20

Ser Phe Gly Asn Thr Ile Thr Cys Tyr Ile Lys Ala Thr Ala Ala Ala 40 35

Arg Ala Ala Gly Leu Arg Asn Pro Asp Phe Leu Val Cys Gly Asp Asp 50

Leu Val Val Val Ala Glu Ser 65

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Ser Phe Gly Asn Thr Ile Thr Cys Tyr Ile Lys A	Ala Thr Ala Ala Ala 45
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Leu Val Val Ala Glu Ser 65 70	·
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Ser Pho	e Gly 35	Asn	Thr	Ile	Thr	Cys 40	Tyr	Ile	Lys	Ala	Thr 45	Ala	Ala	Ala	
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11 213

DNA

<213> hepatitis C virus

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Cys Ser Phe	e Ser Ile	Phe Leu Le 40	ı Ala Leu	Phe Ser	Cys Leu 45	Ile His
Pro Ala Ala 50	Ser Leu	Glu Trp Arg 55	g Asn Thr	Ser Gly 60	Leu Tyr	Val Leu
Thr Asn Asp	-	Asn Ser Se 70	r Ile Val	Tyr Glu 75	Ala Asp	Asp Val 80
Ile Leu His	Thr Pro	Gly Cys Va	l Pro Cys 90	Val Gln	Asp Gly	Asn Thr 95
Ser Ala Cys	Trp Thr	Pro Val Th	r Pro Thr 105	Val Ala	Val Arg 110	Tyr Val
Gly Ala Thi		Ser Ile Ar		Val Asp	Met Leu 125	Val Gly
Ala Ala Thi	Met Cys	Ser Ala Le 135	u Tyr Val	Gly Asp		Gly Ala
Val Phe Let	ı Val Gly	Gln Ala Ph 150	e Thr Phe	Arg Pro	Arg Arg	His Gln 160
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tatgtgtggg	gccgtcttcc	tcgtgggaca	agccttcacg	ttcagacctc	gtcgccatca	480
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<210> 16

<211> 180

<212> PRT

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<400> 16

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Pro Ala Leu Phe Ser Cys Leu Ile His $35 \hspace{1cm} 40 \hspace{1cm} 45$

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

1	le	Leu	His	Thr	Pro 85	Gly	Cys	Val	Pro	Cys 90	Val	Gln	Asp	Gly	Asn 95	Thr		
5	Ser	Ala	Cys	Trp 100	Thr	Pro	Val	Thr	Pro 105	Thr	Val	Ala	Val	Arg 110	Tyr	Val		
C	Bly	Ala	Thr. 115	Thr	Ala	Ser	Ile	Arg 120	Arg	His	Val	Asp	Ile 125	Leu	Val	Gly		
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<212> PRT

<213> hepatitis C virus

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Ala Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val
100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Arg His Val Asp Ile Leu Val Gly 115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly 165 170 175

His Arg Met Ala 180

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<212> PRT

<213> hepatitis C virus

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Ser Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Thr Pro Gly Cys Ile Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Lys Tyr Val

100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly
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His Arg Met Ala 180

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<211> . 541

<212> DNA

<213> hepatitis C virus

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<211> 180

<212> PRT

<213> hepatitis C virus

<400> 22

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<213> hepatitis C virus

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<212> PRT

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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 135 Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 150 155 145 Thr Val Gln Thr Cys Asn Cys Ser Leu Tyr Pro Gly His Leu Ser Gly His Arg Met Ala 180 <210> 25 <211> 541 <212> DNA <213> hepatitis C virus <400> 25 cgtcggcgct cccgtaggag gcgtcgcaag agcccttgcg catggcgtga gggcccttga 60 qqacqqqata aacttcqcaa cagggaattt gcccggttgc tctttttcta tcttccttct 120 tgctctgttc tcttgcttaa tccatccagc agctggtcta gagtggcgga atacgtctgg 180 cctctatgtc cttaccaacg actgttccaa tagtagtatt gtgtatgagg ccgatgacgt 240 tattctgcac gcgcccggct gtgtaccttg tgtccaggac ggcaatacgt ctacatgctg 300 gaccccagta acacctacag tggcagtcag gtacgtcggg gcaaccaccg cttcgatacg 360 cagtcatgtg gacctgttag taggcgcggc cacgatgtgc tctgcgcttt acgtgggtga 420 tatgtgtggg gccgtcttcc tcgtgggaca agccttcacg ttcagacccc gccgccatca 480 aacggtccag acctgtaact gctcgctgta cccaggccat ctttcaggac atcgaatggc 540 541 t <210> 26 <211> 180 <212> PRT hepatitis C virus <213> <400> 26

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Arg Ala Leu Glu Asp Gly Ile Asn Phe Ala Thr Gly Asn Leu Pro Gly 20 25 30

lys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45
Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60
Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 70 75 80
le Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95
Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val 100 105 110
Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly 115 120 125
Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140
Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160
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Cys Ser Phe Ser Ile Phe Leu Leu Ala Leu Phe Ser Cys Leu Ile His 35 40 45

Pro Ala Ala Gly Leu Glu Trp Arg Asn Thr Ser Gly Leu Tyr Val Leu 50 55 60

Thr Asn Asp Cys Ser Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Val 65 70 75 80

Ile Leu His Ala Pro Gly Cys Val Pro Cys Val Gln Asp Gly Asn Thr 85 90 95

Ser Thr Cys Trp Thr Pro Val Thr Pro Thr Val Ala Val Arg Tyr Val
100 105 110

Gly Ala Thr Thr Ala Ser Ile Arg Ser His Val Asp Leu Leu Val Gly
115 120 125

Ala Ala Thr Met Cys Ser Ala Leu Tyr Val Gly Asp Met Cys Gly Ala 130 135 140

Val Phe Leu Val Gly Gln Ala Phe Thr Phe Arg Pro Arg Arg His Gln 145 150 155 160

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Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser

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ctga	atct	gga a	agtaa	accad	cc ag	gcaco	etggg	g tgt	tgct	tgg	aggg	ggtc	ete 9	gcggd	cctag	ļ
cggc	cta	ctg (cttgt	cagt	c gg	gctgo	gttg	g taa	atcgt	ggg	tcat	catco	gag (ctggg	gggca	L
agco	ggc	act o	gtto	ccaga	ac aa	aggag	ggtgt	tgt	atca	aaca	gtad	gate	gag a	atgga	aggagt	:
gcto	gca	agc d	gccc	ccata	ac at	cgaa	acaag	gcto	caggt	taat	agco	ccaco	cag 1	ttcaa	aggaga	L
aaat	cct	tgg a	actgo	etgea	ag co	gagco	caçco	c aac	caaca	aagc	tgt	catto	gag (cccgt	aatag	j
ctto	caa	ctg g	gcaaa	agct	ct ga	aaaco	cttct	gg	cacaa	agca	t					
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Gln 1	Asn	Glu		Cys 5	Leu	Thr	His	Pro	Val 10	Thr	Lys	Tyr	Ile	Met 15	Ala	
Cys	Met	Ser	Ala 20	Asp	Leu	Glu	Val	Thr 25	Thr	Ser	Thr	Trp	Val 30	Leu	Leu	
Gly	Gly	Val 35	Leu	Ala	Ala	Leu	Ala 40	Ala	Tyr	Cys	Leu	Ser 45	Val	Gly	Cys	
Val	Val 50	Ile	Val	Gly	His	Ile 55	Glu	Leu	Gly	Gly	Lys 60	Pro	Ala	Leu	Val	
Pro 65	Asp	Lys	Glu	Val	Leu 70	Tyr	Gln	Gln	Tyr	Asp 75	Glu	Met	Glu	Glu	Cys 80	
Ser	Gln	Ala	Ala	Pro 85	Tyr	Ile	Glu	Gln	Ala 90	Gln	Val	Ile	Ala	His 95	Gln	
Phe	Lys	Glu	Lys 100	Ile	Leu	Gly	Leu	Leu 105	Gln	Arg	Ala	Thr	Gln 110	Gln	Gln	

Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln Lys Leu Glu Thr

115 120 125

Phe Trp His Lys His 130

<210> 33

<211> 401

<212> DNA

<213> hepatitis C virus

<400> 33

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cggcctactg cttgtcagtc ggctgcgttg taatcgtggg tcatatcgag ctggggggca 180
agccggcact cgttccagac aaggaggtgt tgtatcaaca gtacgatgag atggaggagt 240
gctcgcaagc cgccccatac atcgaacaag ctcaggtaat agcccaccag ttcaaggaga 300
aaatccttgg actgctgcag cgagccaccc aacaacaagc tgtcattgag cccgtaatag 360
cttccaactg gcaaaagctt gaaaccttct ggcacaagca t 401

<210> 34

<211> 133

<212> PRT

<213> hepatitis C virus

<400> 34

Gln Asn Glu Ile Cys Leu Thr His Pro Val Thr Lys Tyr Ile Met Ala 1 5 10 15

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu 20 25 30

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Leu Val 50 55 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Ile Leu Gly Leu Leu Gln Arg Ala 100 105	Thr Gln Gln Gln 110
Ala Val Ile Glu Pro Val Ile Ala Ser Asn Trp Gln 115 120	Lys Leu Glu Thr 125
Phe Trp His Lys His	
<210> 35 <211> 401 <212> DNA <213> hepatitis C virus	
<400> 35	
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ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggg	gteete geggeeetag 120
cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcat	atcgag ctggggggca 180
agccggcaat cgttccagac aaagaggtgt tgtatcaaca atac	gatgag atggaagagt 240
gctcacaage tgccccatat atcgaacaag ctcaggtaat agct	caccag ttcaagggaa 300
aagteettgg attgetgeag egageeacce aacaacaage tgte	attgag cccatagtaa 360
ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t	401
<210> 36 <211> 133 <212> PRT <213> hepatitis C virus	
<400> 36	
Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys 1 10	Tyr Ile Met Ala 15
Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr	Trp Val Leu Leu 30
Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu	Ser Val Gly Cys

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val

55

. 35

50

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Gly Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 105 100 Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 37 <211> 401 <212> DNA <213> hepatitis C virus <400> 37 tccaaaatga aatctgcttg acacacccca tcacaaaata catcatggca tgcatgtcag 60 ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 180 cggcctactg cttgtcagtc ggttgtgttg tgattgtggg tcatatcgag ctggggggca agccggcaat cgttccagac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 gctcacaagc tgccccatat atcgaacaag ctcaggtgat agctcaccag ttcaaggaaa 300 aagtccttgg attgctgcag cgagccaccc aacaacaagc tgtcattgag cccatagtaa 360 401 ctaccaactg gcaaáagctt gaggcctttt ggcacaagca t <210> 38 <211> - 133 <212> PRT <213> hepatitis C virus <400> Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala 10 5

Cys Met Ser Ala Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 40 Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 55 Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 100. Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 120 115 Phe Trp His Lys His 130 <210> 39 <211> 401 <212> DNA <213> hepatitis C virus <400> 60 tccaaaatga aatctgcttg acacacccca tcacaaaata catcatggca tgcatgtcag ctgatctgga agtaaccacc agcacctggg ttttgcttgg aggggtcctc gcggccctag 120 concetacte cttetcagte egttetette teattetege teatategag ctegggggea 180 aqccqqcaat cqttccaqac aaagaggtgt tgtatcaaca atacgatgag atggaagagt 240 qctcacaaqc tqccccatat atcgaacaag ctcaggtaat agctcaccag ttcaaggaaa 300 360 aaqtccttqq attqctqcaq cqaqccaccc aacaacaagc tgtcattgag cccatagtaa

<210> 40

<211> 133

<212> PRT

<213> hepatitis C virus

ctaccaactg gcaaaagctt gaggcctttt ggcacaagca t

<400> 40

Gln Asn Glu Ile Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala

401

5

Cys	Met	Ser	Ala	Asp	Leu	Glu	Val	Thr	Thr	Ser	Thr	Trp	Val	Leu	Leu
			20					25					30		

10

Gly Gly Val Leu Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys 35 40 45

Val Val Ile Val Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val 50 60

Pro Asp Lys Glu Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys 70 75 80

Ser Gln Ala Ala Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln 85 90 95

Phe Lys Glu Lys Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln 100 105 110

Ala Val Ile Glu Pro Ile Val Thr Thr Asn Trp Gln Lys Leu Glu Ala 115 120 125

Phe Trp His Lys His

<210> 41

<211> 509

<212> DNA

<213> hepatitis C virus

<400> 41

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<210> 42

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 42

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 43

<211> 509

<212> DNA

<213> hepatitis C virus

-100- 12						
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ggggccctag	gatgggtgtg	cgcgcgactc	ggaagacttc	ggaacggtcg	caaccccgtg	180
gacggcgtca	gcctattccc	aaggcgcgcc	agcccacggg	ccggtcctgg	ggtcaacccg	240
ggtacccttg	gcccctttac	gccaatgagg	gcctcgggtg	ggcagggtgg	ctgctctccc	300
ctcgaggctc	tcggcctaat	tggggcccca	atgacccccg	gcgaaaatcg	cgtaatttgg	360
gtaaggtcat	cgatacccta	acgtgcggat	tcgccgatct	catggggtat	atcccgctcg	420
taggcggccc	cattgggggc	gtcgcaaggg	ctctcgcaca	cggtgtgagg	gtccttgagg	480
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<210> 44

<211> 169

<212> PRT

<213> hepatitis C virus

<400> 44

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Ile 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Ala Thr Gly Asn Leu 165

<210> 45

<211> 580

<212> DNA

<213> hepatitis C virus

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<210> 46

<211> 193

<212> PRT

<213> hepatitis C virus

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<400> 46

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

ggtaggccaa atgttcacct ataggcctcg ccagcacgct acggtgcaga actgcaactg

540

580

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 55 Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 70 Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys 105 100 Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120 Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 150 145 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met 185 180 Ala <210> 47 580 <211> <212> DNA <213> hepatitis C virus <400> 47 aacgtgcgga ttcgccgatc tcatggggta tatcccgctc gtaggcggcc ccattggggg 60 cgtcgcaagg gctctcgcac acggtgtgag ggtccttgag gacggggtaa actatgcaac 120 agggaattta cccggttgct ctttctctat ctttattctt gctcttctct cgtgtctgac 180

cgttccggcc	tctgcagttc	cctaccgaaa	tgcctctggg	atttatcatg	ttaccaatga	240
ttgcccaaac	tcttccatag	tctatgaggc	agataacctg	atcctacacg	cacctggttg	300
cgtgccttgt	gtcatgacag	gtaatgtgag	tagatgctgg	gtccaaatta	cccctacact	360
gtcagccccg	agcctcggag	cagtcacggc	tactattagg	agagccgttg	actacctagc	420
gggaggggct	gccctctgct	ccgcgttata	cgtaggagac	gcgtgtgggg	cactattctt	480
ggtaggccaa	atgttcacct	ataggcctcg	ccagcacgct	acggtgcaga	actgcaactg	540
ttccatttac	agtggccatg	ttaccggcca	ccggatggca	•		580

<210> 48

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 48

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Ile Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu
20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 115 120 125

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 130 135 140

Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 165 170 175

Asn Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210> 49

<211> 959

<212> DNA

<213> hepatitis C virus

<400> 49

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<212> PRT

<213> hepatitis C virus

<400> 50

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro
100 105 110

Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175

Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205 Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro 210 215 220

Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val 225 230 235 240

Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255

Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys 260 265 270

Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly 275 280 285

Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 290 295 300

Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala 305 310 315

<210> 51

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tectacaege acetggttge gtgeettgtg teatgacagg taatgtgagt agatgetggg							
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gageegttga etacetageg ggaggggetg ecetetgete egegttatae gtaggagaeg							
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<210> 52 <211> 319 <212> PRT							
<213> hepatitis C virus							
<400> 52							
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Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30							
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Met Gly Val Arg Ala 35 40 45							
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60							
Ile Pro Lys Ala Arg Gln Pro Thr Gly Arg Ser Trp Gly Gln Pro Gly 65 70 75 80							
Tyr Pro Trp Pro Leu Tyr Ala Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95							
Leu Leu Ser Pro Arg Gly Ser Arg Pro Asn Trp Gly Pro Asn Asp Pro 100 105 110							
Arg Arg Lys Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys 115 120 125							

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Gly Pro Val

•• 1		
e.		
	Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile	
	165 170 175	
	Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val	
	180 185 190	
	and the state of t	
	Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp Cys Pro 195 200 205	
	Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His Ala Pro	
	210 215 220	
		_
	Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys Trp Val	
	225 230 235 240	
	Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val Thr Ala 245 250 255	
	Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala Leu Cys	
	260 265 270	
	Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu Val Gly	
	275 280 285	
	Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln Asn Cys 290 295 300	
	Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Arg Met Ala	
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	gacggcgtca gcctattccc aaggcgcgcc agcccacggg ccggtcctgg ggtcaacccg 240	
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Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser Ala Val 180 185 190

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120

180

240

300

354

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Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala Thr Xaa

Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val

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Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Xaa Thr Xaa 50 55 60

Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu Glu Val 65 70 75 80

Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Xaa Arg Ala Ile Pro 85 90 95

Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His Ser Lys
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Glu Cys Ser Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala 65 70 75 80

Gly Gln Phe Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln 85 90 95

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Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Lys Ala 65 70 75 80

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Ala

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catgcataac agcaggggag acctgtgcgg gtatcgcaga tgccgtgcga gcggcgtcta 180
caccaccagc ttcgggaaca cactgacgtg ctatctcaaa gcctcagccg ctatcagagc 240
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Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Arg Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ala Gly Leu Arg Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
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Ala

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catgcataat agcaaaggag acctgtgcgg gtatcgcaga tgccgcgcaa gcggcgtcta 180
caccaccagc ttcgggaaca cactgacgtg ctatctcaaa gcctcagccg ccatcagggc 240
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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

Ser Gly Leu Arg Asp Cys Thr Met Leu Val Tyr Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Asp Gly Val Glu Glu Asp Lys Arg Ala Leu Gly
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	ggcggggctg	agagactgca	ccatgttggt	ctgtggtgat	gacctggtcg	tcatcgctga	300
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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Ala Ile Thr Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met His Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Ser Ala Ala Ile Arg Ala 65 70 75 80

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Cys Cys Asp Leu Glu Pro Glu Ala Arg Lys Val Ile Ser Ala Leu Thr 20 25 30

Glu Arg Leu Tyr Val Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Val Ala Ala Thr Arg Ala 65 70 75 80

Ala Gly Leu Lys Gly Cys Ser Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Cys Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 100 105 110

Ala

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Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Lys Arg Ala Leu Gly

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120

180

240

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340

Ala

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Pro	Val	Gly	Gly 20	Val	Ala	Arg	Ala	Leu 25	Ala	His	Gly	Val	Arg 30	Ala	Val
Glu	Asp	Gly 35	Ile	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Cys	Ser	Phe
Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Thr	Ser
Ala 65	Val	Asn	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thr	Asn	Asp 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Thr Glu His His Ile Leu His 85 90 95

Leu Pro Gly Cys Leu Pro Cys Val Arg Val Gly Asn Gln Ser Arg Cys Trp Val Ala Leu Thr Pro Thr Val Ala Ala Pro Tyr Ile Gly Ala Pro 115 Leu Glu Ser Leu Arg Ser His Val Asp Leu Met Val Gly Ala Ala Thr 135 130 Ala Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 155 150 145 Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 170 165 Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg 180 <210> 120 <211> 574 <212> DNA hepatitis C virus <213> <400> 120 acgtgcggct ttgccgacct catgggatac atcccgctcg tgggcgcccc tgtgggtggc 60 gtcgccaggg ccttggcaca tggtgtcagg gccgtggagg acgggattaa ctatgcaaca 120 gggaatette ceggttgete ettttetate tteettetag caettetete gtgettgaet 180 gtcccggcct cggcgcagca ctaccggaac atctcgggca tttatcacgt caccaatgac 240 300 tgcccqaact ctaqtataqt qtatgaagct gaccatcata tcatgcatct accagggtgt gtgccttgcg tgagaaccgg gaacacctcg cgctgctggg ttcctttaac acccactgtg 360 gctgccccct atgttggcgc gccgctcgaa tccatgcggc ggcacgtgga cttaatggtg 420 ggtgccgcca ccgtctgctc ggccctgtac atcggagacc tttgcggagg tgtcttcctg 480 gtcgggcaga tgttcacctt ccggccgcgc cgccattgga ctacccagga ctgcaactgc 540 574 tctatctatg atggccacat caccggccat agaa

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Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ile Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Thr Ser Arg Cys
100 105 110

Trp Val Pro Leu Thr Pro Thr Val Ala Ala Pro Tyr Val Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Arg Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Asp Gly His Ile Thr Gly His Arg 180 185 190

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ggtgcggcca	ccgtgtgttc	tgtcctctat	gttggagacc	tctgtggagg	tgctttccta	480
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Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Glu His Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Val Val Tyr Glu Thr Asp His His Ile Leu His
85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Ala Gly Asn Val Ser Arg Cys
100 105 110

Trp Thr Pro Val Thr Pro Thr Val Ala Ala Val Ser Met Asp Ala Pro 115 120 125

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Val Gl	y Gln	Met	Phe 165	Thr	Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln		
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Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile

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319 <211>

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- Arg Arg Pro Gln Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30
- Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45
- Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60
- Ile Pro Lys Asp Arg Arg Pro Thr Gly Lys Ser Trp Gly Lys Pro Gly 65 70 75 80
- Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Leu Gly Trp Ala Gly Trp 85 90 95
- Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Thr Asp Pro 100 105 110
- Arg His Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
 115 120 125
- Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Val Val Gly Ala Pro Val 130 135 140
- Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu Glu Asp 145 150 155 160
- Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe Ser Ile 165 170 175
- Phe Leu Leu Ala Leu Leu Ser Cys Ile Thr Val Pro Val Ser Gly Leu 180 185 190
- Gln Val Lys Asn Thr Ser Ser Ser Tyr Met Val Thr Asn Asp Cys Gln 195 200 205
- Asn Ser Ser Ile Val Trp Gln Leu Arg Asp Ala Val Leu His Val Pro 210 215 220
- Gly Cys Val Pro Cys Glu Glu Lys Gly Asn Ile Ser Arg Cys Trp Ile

Pro Val Ser Pr	o Asn Ile Ala 245	a Val Ser Gl 25		Ala Leu Thr 255	Lys								
Gly Leu Arg Th		o Thr Ile II 265	ie Ala Ser	Ala Thr Phe 270	Cys								
Ser Ala Leu Ty 275	r Ile Gly As	Leu Cys Gl	ly Ala Val	Met Leu Ala 285	Ser								
Gln Val Phe Il 290	le Ile Ser Pro 29		is Lys Phe 300	Val Gln Asp	Cys								
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catgctaaac ago	caaagggc aaac	ctgcgg atac	agacgc tgcc	gegeca gegg	agtgtt 18								
caccactage at	gggaaata ccat	cacgtg ctac	gtgaag gcac	aagcag cctg	taaggc 24								
tgcgggcata at	tgccccca cgat	gctggt gtgc	ggcgac gato	tagttg tcat	ctcaga 30								
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<400> 146

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Cys	Gly 50	Tyr	Arg	Arg	CAa	Arg 55	Ala	Ser	Gly	Val	Phe 60	Thr	Thr	Ser	Met	
Gly 65	Asn	Thr	Ile	Thr	Cys 70	Tyr	Val	Lys	Ala	Gln 75	Ala	Ala	Cys	Lys	Ala 80	
Ala	Gly	Ile	Ile	Ala 85	Pro	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
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Ala									٠					•		
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gcc	ccca	gtt	gggt	gtgc	gt g	cagt	gcgc	a ag	actt	ccga	gcg	gtcg	caa	cctc	gcagta	180
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acc	cttg	gcc	ccta	tatg	gg a	atga	gggc	t gc	gġgt	gggc	agg	gtgg	ctc	ctgt	ccccgc	300
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30

Glu Phe Thr Cys Tyr His Ala Gly Ala Pro Ser Trp Val Cys Val Gln Cys Ala Arg Leu Pro Ser Gly Arg Asn Leu Ala Val Gly Ala Asn Pro 55 Ser Pro Gly Arg Ala Glu Pro Arg Ala Gly Pro Gly Leu Ser Pro Gly 70 Thr Leu Gly Pro Tyr Met Gly Met Arg Ala Ala Gly Gly Gln Gly Gly 90 85 Ser Cys Pro Arg Ala Ala Leu Ala Arg Arg Gly Ala Gln Met Thr Pro 105 Gly Ala Gly 115 <210> 149 <211> 280 <212> DNA <213> hepatitis C virus <400> 149 ggcctgtgac ctcaaggacg aggctaggag ggtgataact tcactcacgg agcggcttta 60 ctgtggtggt cctatgttca acagcaaggg acaacactgc ggttaccgcc gctgccgtgc. 120 tagtggggtg ctacccacca gcttcgggaa cacaatcacc tgttacatca aagcaaaggc 180 agctaccaaa gctgccggaa ttaaaaatcc atcattcctt gtctgcggag atgacttggt 240 280 cgtgattgct gagagtgcag ggatcgatga ggacagagcg <210> 150 <211> 93 <212> PRT <213> hepatitis C virus 150 <400> Ala Cys Asp Leu Lys Asp Glu Ala Arg Arg Val Ile Thr Ser Leu Thr 10 Glu Arg Leu Tyr Cys Gly Gly Pro Met Phe Asn Ser Lys Gly Gln His

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Thr Arg Ly	s Thr Se	er Glu Ar 55		Gln	Pro	Arg	Gly 60	Arg	Arg	Gln	Pro	
Ile Pro Ly 65	rs Ala Ar	g Gln Pr 70	o Thr	Gly .		Ser 75	Trp	Gly	Gln	Pro	Gly 80	
Tyr Pro Tr	p Pro Le 85		a Asn		Gly 90	Leu	Gly	Trp	Ala	Gly 95	Trp	
Leu Leu Se	er Pro Ar 100	cg Gly Se	r Arg	Pro 105	Asn	Trp	Gly	Pro	Asn 110	Asp	Pro	
Arg Arg Ly		rg Asn Le	u Gly 120	Lys	Val	Ile	Asp	Thr 125	Leu	Thr	Cys	
Gly Phe Al	la Asp Le		y Tyr 5	Ile	Pro	Leu	Val 140	Gly	Gly	Pro	Ile	
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Gly Val As	_	la Thr 65					•					
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gtgccttgt	g tcatga	cagg taa	tgtgag	rt aga	atgc	tggg	tcc	aaat	tac	ccct	acactg	36
tcagccccg	a gcctcg	gagc agt	cacggc	t cci	tctt	cgga	gag	ccgt	tga	ctac	ctagcg	42

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<211> 193

<212> PRT

<213> hepatitis C virus

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Glu Asp Gly Val Asn Tyr Pro Thr Gly Asn Leu Pro Gly Cys Ser Phe 40 35

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 55 50

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 75 70 65

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asn Leu Ile Leu His 85

Ala Pro Gly Cys Val Pro Cys Val Met Thr Gly Asn Val Ser Arg Cys

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Leu Gly Ala Val 120

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Ala Gly Gly Ala Ala 135 130

Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 155 150 145

Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln . 170 175 165

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Ala

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gtaggco	caaa	tgttcaccta	taggcctcgc	cagcatgcta	cggtgcagga	ctgcaactgt	540
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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 156

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Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 70 Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 150 145 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 170 165 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 180 Ala <210> 157 530 <211> DNA <212> hepatitis C virus <213> <400> 157 cacctacgac agetetgetg gtggcccagt tactgcggat tececaagtg gtcattgaca 60 tcatcgcagg gagccactgg ggggtcttgt ttgccgccgc atactatgca tcggtggcta 120 actggaccaa ggtcgtgctg gtcttgtttc tgtttgcagg ggttgatgct actacccaga 180 tttcgggcgg ctccagcgcc caaacgacgt atggcatcgc ctcatttatc acccgcggcg 240 cgcagcagaa actgcagctc ataaatacca acggaagctg gcacatcaac aggaccgccc 300 ttaattgtaa tgacagcctc cagactgggt tcatagccgg cctcttctac taccataagt 360 tcaactcttc tggatgcccg gatcggatgg ctagctgtag ggcccttgcc acttttgacc 420 agggctgggg aactatcagc tatgccaaca tatcgggtcc cagtgatgac aaaccatatt 480 <210> 158

<211> 176

<212> PRT

<213> hepatitis C virus

<400> 158

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Val Ile Asp Ile Ile Ala Gly Ser His Trp Gly Val Leu Phe Ala Ala 25 20

Ala Tyr Tyr Ala Ser Val Ala Asn Trp Thr Lys Val Val Leu Val Leu 35

Phe Leu Phe Ala Gly Val Asp Ala Thr Thr Gln Ile Ser Gly Gly Ser 50

Ser Ala Gln Thr Thr Tyr Gly Ile Ala Ser Phe Ile Thr Arg Gly Ala 70

Gln Gln Lys Leu Gln Leu Ile Asn Thr Asn Gly Ser Trp His Ile Asn

Arg Thr Ala Leu Asn Cys Asn Asp Ser Leu Gln Thr Gly Phe Ile Ala 105 100

Gly Leu Phe Tyr Tyr His Lys Phe Asn Ser Ser Gly Cys Pro Asp Arg 120 115

Met Ala Ser Cys Arg Ala Leu Ala Thr Phe Asp Gln Gly Trp Gly Thr 135 130

Ile Ser Tyr Ala Asn Ile Ser Gly Pro Ser Asp Asp Lys Pro Tyr Cys 155 145 150

Trp His Tyr Pro Pro Arg Pro Cys Gly Val Val Pro Ala Gln Glu Val 170

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Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Ser Leu Arg

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120

180

240

300

340

Ala

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catgtataac agcaaggggc aacaatgtgg ttatcgcaga tgccgcgcca gcggcgtctt
caccaccagt atgggcaaca ccatgacgtg ctacattaag gctttagcct cctgtagaac
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Ser Cys Asp Ser Gln Pro Asp Ala Arg Ala Ala Ile Arg Ser Leu Thr 20 25 30
Gln Arg Leu Phe Cys Gly Gly Pro Met Tyr Asn Ser Lys Gly Gln Gln 35 40 45
Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60
Gly Asn Thr Met Thr Cys Tyr Ile Lys Ala Leu Ala Ser Cys Arg Thr 65 70 75 80

Ala Ile Cys Glu Ser Gln Gly Thr His Glu Asp Glu Ala Asn Leu Arg 100 105 110

Ala Gly Leu Arg Asp Tyr Thr Leu Leu Val Cys Gly Asp Asp His Val

Ala

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gacgttaagt tcccgggtgg tggcca	agatc gtt	ggcggag	tttacttgt	t gccgcgcagg
			agggtggg	2 2001001000
ggccctaggt tgggtgtgcg cgcgac	ercgg aag	Jacttegg	ageggeege	a accregigg
aggegecaae etateeceaa ggegeg	gccga tcc	gagggca	gatcctggg	c gcagcccggg
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cgcgggtctc ggccgtcttg gggcc	ctaat gat	ccccggc	ggaggtccc	g caacctgggt
•				
aaggtcatcg ataccctaac atgcg	gette ged	gaeetca	Lygyataca	t coogeologia
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010 164				
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Ara Ara Dro Met Ash Val Ivs	Phe Pro	Glv Glv	Glv Gln 1	le Val Gly
Arg Arg Pro Met Asp Val Lys	Phe Pro 25	Gly Gly		Ile Val Gly
Arg Arg Pro Met Asp Val Lys 20		Gly Gly		
20	25		3	
20 Gly Val Tyr Leu Leu Pro Arg	25 Arg Gly		Leu Gly V	
20	25		3	
20 Gly Val Tyr Leu Leu Pro Arg	25 Arg Gly		Leu Gly V	
20 Gly Val Tyr Leu Leu Pro Arg	25 Arg Gly 40	Pro Arg	Leu Gly V 45	0 Val Arg Ala
20 Gly Val Tyr Leu Leu Pro Arg 35	25 Arg Gly 40	Pro Arg	Leu Gly V	0 Val Arg Ala
Gly Val Tyr Leu Leu Pro Arg 35	25 Arg Gly 40	Pro Arg	Leu Gly V 45	0 Val Arg Ala
Gly Val Tyr Leu Leu Pro Arg 35 Thr Arg Lys Thr Ser Glu Arg 50 55	25 Arg Gly 40 Ser Gln	Pro Arg	Leu Gly V 45 Gly Arg A	Val Arg Ala Arg Gln Pro
Gly Val Tyr Leu Leu Pro Arg 35	25 Arg Gly 40 Ser Gln	Pro Arg	Leu Gly V 45 Gly Arg A	Val Arg Ala Arg Gln Pro
Gly Val Tyr Leu Leu Pro Arg 35 Thr Arg Lys Thr Ser Glu Arg 50 Ile Pro Lys Ala Arg Arg Ser	25 Arg Gly 40 Ser Gln	Pro Arg Pro Arg	Leu Gly V 45 Gly Arg A	Val Arg Ala Arg Gln Pro Gln Pro Gly
Gly Val Tyr Leu Leu Pro Arg 35 Thr Arg Lys Thr Ser Glu Arg 50 Ile Pro Lys Ala Arg Arg Ser	25 Arg Gly 40 Ser Gln Glu Gly	Pro Arg Pro Arg Arg Ser	Leu Gly V 45 Gly Arg A	Val Arg Ala Arg Gln Pro Gln Pro Gly 80

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro

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<213> hepatitis C virus

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<222> (114)..(114)

<223> "Xaa" is any amino acid

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Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Ala

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Thr Glu Gly Arg Ser Trp Ala Gln Pro Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Xaa Ser Arg Xaa Ser Trp Gly Pro Asn Asp Pro 100 105 110

Arg Xaa Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu 115 120 125

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<211> 193

<212> PRT

<213> hepatitis C virus

<400> 168

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln
165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<210> 169

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<212> DNA

<213> hepatitis C virus

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<212> PRT

<213> hepatitis C virus

<400> 170

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Gly Val Asn Tyr Arg Asn Ala Ser Gly Val Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Asp Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Val Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Val Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Ile Thr Gly His Arg Met 180 185 190

<400> 172

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gtccccg	gcct	cggctgtgca	ttatcacaac	acctcgggca	tctaccacct	caccaatgac	240
tgcccta	act	ctagcatagt	ctttgaggca	gtccatcaca	tcttgcacct	tccaggatgc	300
gtccctt	gtg	taagaactgg	gaaccagtct	cggtgctggg	tagccttgac	ccccacgctg	360
gccgcgc	ccat	acctiggcgc	tccactcgag	tccatgcggc	gtcacgtgga	tttgatggtg	420
ggcacto	gcta	cattgtgctc	agcactctac	gttggggacc	tgtgcggggg	catattccta	480
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Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Leu Thr Asn Asp

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Val His His Ile Leu His 90 85

70

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys 105 100

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro Tyr Leu Gly Ala Pro 120 115

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr 140 130

Leu Cys Ser Ala Leu Tyr Val Gly Asp Leu Cys Gly Gly Ile Phe Leu 150

Ala Gly Gln Met Phe Thr Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165

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Ala

<210> 173

<211> 579

<212> DNA

<213> hepatitis C virus

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<210> 174

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 174

Thr Cys Gly Ser Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val His Tyr His Asn Thr Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Phe Glu Ala Glu His His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Ile Ala Leu Thr Pro Thr Leu Ala Ala Pro His Ile Gly Ala Pro 115 120 125

Leu Glu Ser Met Arg Arg His Val Asp Leu Met Val Gly Thr Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Ile Phe Leu 145 150 155 160

Val Gly Gln Met Phe Asn Phe Arg Pro Arg Leu His Trp Thr Thr Gln 165 170 175

Glu Cys Asn Cys Ser Ile Tyr Pro Gly His Ile Thr Gly His Arg Met

180 185 190

Ala

	175 579 DNA				•		
		titis C vir	rus				
	175 gct	ttgccgacct	catgggatac	atcccgctcg	tgggcgcccc	tgtgggtggc	60
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gtcccgg	gcct	cggcgcagca	ctaccggaac	atctcgggca	tttatcacgt	caccaatgac	240
tgcccga	act	ctagtatagt	gtatgaagct	gaccatcata	tcatgcatct	accagggtgt	300
gtgcctt	gcg	tgagaaccgg	gaacacctcg	cgctgctggg	ttcctttaac	acccactgtg	360
gctgccc	cct	atgttggcgc	gccgctcgaa	tccatgcggc	ggcacgtgga	cttaatggtg	420
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<210> 176

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 176

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala 65	Gln	His	Tyr	Arg	Asn 70	Ile	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80	
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	His	His	Ile	Met 95	His	
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Thr	Gly	Asn	Thr	Ser 110	Arg	Cys	
Trp	Val	Pro 115	Leu	Thr	Pro	Thr	Val. 120	Ala	Ala	Pro	Tyr	Val 125	Gly	Ala	Pro	:
Leu	Glu 130	Ser	Met	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr	
Val 145		Ser	Ala	Leu	Tyr 150	Ile	Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160	
Val	Gly	Gln	Met	Phe 165	Thr	Phe	Arg	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln	
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Ala			,													
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gg 9	aacc	ttc	ccgg	ctgc	tc c	tttt	ctat	c tt	cctc	ttgg	tac	ttct	ctc	gcgc	ctaact	180
gto	ccag	cgt	ctgc	tcag	ca c	tacc	ggaa	t gc	atcg	ggca	tct	acca	tgt	cacc	aacgac	240
tgo	ccga	act	ccag	tatt	gt g	tatg	aagc	c ga	ccat	caca	tca	tgca	cct	accc	gggtgt	300
gtg	rccct	gtg	taag	aact	gg g	aatg	tctc	g cg	ttgc	tgga	ttc	cttt	aac	accc	actgta	360
gcc	gtcc	cct	acct	.cggg	gc t	ccac	ttac	g tc	tgta	cggc	agc	atgt	gga	cctg	atggtg	420
999	ıgcgg	cca	cctt	atgo	tc t	gccc	tcta	c at	cgga	gacc	att	gcgg	agg	tgtc	ttcttg	480

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579

<210> 178

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 178

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Val Leu Leu Ser Arg Leu Thr Val Pro Ala Ser 50 55 60

Ala Gln His Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp His His Ile Met His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Val Ser Arg Cys
100 105 110

Trp Ile Pro Leu Thr Pro Thr Val Ala Val Pro Tyr Leu Gly Ala Pro 115 120 125

Leu Thr Ser Val Arg Gln His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Leu Cys Ser Ala Leu Tyr Ile Gly Asp His Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Met Val Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln
165 170 175

Asp Cys Asn Cys Ser Ile Tyr Val Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<223>

"Xaa" is any amino acid

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<210>
     179
<211> 579
<212> DNA
      hepatitis C virus
<213>
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<221> misc feature
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      (568)..(579)
<223> "n" is any nucleotide
<400>. 179
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                                                                      120
gggaatette etggttgete tttetetate tecettettg aacttetete gtgeetgaet
                                                                      180
gttcccgcct cagccatcca ctatcgcaat gcttcggacg gttattatat caccaatgat
                                                                      240
tgcccgaact ctagcatagt gtatgaagcc gagaaccaca tcttgcacct tccggggtgt
                                                                      300
ataccetgtg tgaagacegg gaateagteg eggtgetggg tggeteteae eeceaegetg
                                                                      360
geggeeceae acctaegtge teegettteg teettaeggg egeatgtgga cetaatggtg
                                                                      420
ggggccgcca cggcatgctc cgctttttac attggagatc tgtgcggggg tgtgtttttg
                                                                      480
gcgggccaac tgttcactat ccggccacgc attcatgaaa ccactcagga ctgcaattgc
                                                                      540
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tccatctact cagggcacat cacgggtnnn nnnnnnnn
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       "Xaa" is any amino acid
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       MISC FEATURE
<221>
       (190)..(193)
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<400> 180

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Xaa Leu Ala His Gly Val Arg Ala Leu 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Ser Leu Leu Glu Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Ile His Tyr Arg Asn Ala Ser Asp Gly Tyr Tyr Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Ile Pro Cys Val Lys Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Leu Ala Ala Pro His Leu Arg Ala Pro 115 120 125

Leu Ser Ser Leu Arg Ala His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Ala Cys Ser Ala Phe Tyr Ile Gly Asp Leu Cys Gly Gly Val Phe Leu 145 150 155 160

Ala Gly Gln Leu Phe Thr Ile Arg Pro Arg Ile His Glu Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Ile Thr Gly Xaa Xaa Xaa 180 185 190

Xaa

<210> 181 <211> 579

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<213> hepatitis C virus
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                                                                      120
gggaatette etggttgete tttetetate tneettetgg caettetete gtgeetgaet
                                                                      180
gtcccggcct cggctcagca ctaccggaat gtctcgggca tctaccacgt caccaatgat
                                                                      240
tgcccgaatt ccagcatagt gtatgaagcc gatcaccaca tcatgcactt accagggtgc
                                                                      300
ataccetgeg tgaggacegg gaacgttteg egetgetggg tatetetgae acetaetgtg
                                                                      360
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                                                                      420
ggtgcagcca ccctttgctc tgccctctac gtcggagacc tctgtggagg tgtcttccta
                                                                      480
gtgggacaga tgttcacctt ccagccgcgc cgccactgga ccactcagga ctgcaactgc
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DNA

<221> MISC FEATURE

(51)..(51) <222>

"Xaa" is any amino acid <223>

<400> 182

Ala Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35

Ser Ile Xaa Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser

Ala 65	Gln	His	Tyr	Arg	Asn 70	Val	Ser	Gly	Ile	Tyr 75	His	Val	Thr	Asn	Asp 80		
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Ala 90	Asp	His	His	Ile	Met 95	His		
Leu	Pro	Gly	Cys	Ile	Pro	Cys	Val	Arg 105	Thr	Gly	Asn	Val	Ser	Arg	Cys		
Trp	Val	Ser	Leu	Thr	Pro	Thr.	Val 120	Ala	Ala	Pro	Tyr	Leu 125	Gly	Ala	Pro		
Leu	Thr 130	Ser	Leu	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr		
Leu 145	CÀa	Ser	Ala	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160		
Val	Gly	Gln	Met	Phe 165		Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln		
Asp	Cys	Asn	Суs 180		Ile	Tyr	Val	Gly 185	His	Ile	Thr	Gly	His 190		Met		
Ala																	
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gtt	ccca	.cct	cago	cgto	aa c	tato	gcaa	ic go	ctcg	ggcg	tct	atca	tat	cacc	aatga	С	24
tgc	ccga	att	cgag	gcata	ıgt g	tace	gaggo	t ga	ctac	caca	tcc	taca	cct	ccct	gggtg	С	3 0

ttaccctgcg tgagggttgg gaatcagtca cgctgctggg tggcccttac tcccaccgtg

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gttggtcaga	tgttttcttt	ccagccgcga	cgccactgga	ccacgcagga	ctgcaattgt	540
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<210> 184						

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 184

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Leu Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Thr Ser 50 55 60

Ala Val Asn Tyr Arg Asn Ala Ser Gly Ile Tyr His Ile Thr Asn Asp 65 70 75 80

Cys Pro Asn Ala Ser Ile Val Tyr Glu Thr Glu Asn His Ile Leu His 85 90 95

Leu Pro Gly Cys Val Pro Cys Val Arg Thr Gly Asn Gln Ser Arg Cys
100 105 110

Trp Val Ala Leu Thr Pro Thr Val Ala Ser Pro Tyr Ala Gly Ala Pro 115 120 125

Leu Glu Pro Leu Arg Arg His Val Asp Leu Met Val Gly Ala Ala Thr 130 135 140

Met Cys Ser Ala Leu Tyr Ile Gly Asp Leu Cys Gly Gly Leu Phe Leu 145 150 155 160

Val Gly Gln Met Phe Thr Phe Gln Pro Arg Arg His Trp Thr Thr Gln

165 170 175

Asp Cys Asn Cys Ser Ile Tyr Thr Gly His Ile Thr Gly His Arg Met 180 185 190

Ala

<212>	185 579 DNA hepa	ititis C vir	rus				
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gggaato	ctcc	ccggttgctc	tttctctatc	tacctcttgg	cacttctctc	gtgcctgact	180
gttccca	acct	cggccatcca	ctatcgcaat	gcctcgggcg	tctaccacgt	caccaatgac	240
tgcccga	act	cgagcatagt	gtacgaggcc	gaccaccaca	tcctacacct	tccagggtgc	300
ttaccct	gtg	tgagggttgg	gaatcagtca	cgttgttggg	tggccctctc	tcccaccgtg	360
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gttggto	caga	tgttttcttt	ccggccacga	cgccactgga	ctacgcagga	ctgcaattgt	540
tccatct	acg	cggggcacat	cactggccac	ggaatggca			579

<210> 186

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 186

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala 1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Glu His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

		·			
	Ser Ile Tyr 50	Leu Leu Ala Leu Leu 55		Thr Val Pro Thr Ser 50	
	Ala Ile His	s Tyr Arg Asn Ala Ser (Gly Val Tyr 1 75	His Val Thr Asn Asp 80	
	Cys Pro Asr	n Ser Ser Ile Val Tyr 85	Glu Ala Asp 1 90	His His Ile Leu His 95	
:	Leu Pro Gly	y Cys Leu Pro Cys Val 100	Arg Val Gly 1	Asn Gln Ser Arg Cys 110	·
	Trp Val Ala	a Leu Ser Pro Thr Val 5. 120	Ala Ala Pro '	Tyr Ile Gly Ala Pro 125	
	Val Glu Ser 130	r Phe Arg Arg His Val 135		Val Gly Ala Ala Thr 140	
	Val Cys Sei 145	r Ala Leu Tyr Ile Gly 150	Asp Leu Cys 155	Gly Gly Val Phe Leu 160	
	Val Gly Gl	n Met Phe Ser Phe Arg 165	Pro Arg Arg 170	His Trp Thr Thr Gln 175	
	Asp Cys Ası	n Cys Ser Ile Tyr Ala 180	Gly His Ile 185	Thr Gly His Gly Met 190	
	Ala				
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	gggaatcttc	ccggttgctc tttctctatc	c tteetettgg	cacttettte gtgeetgaet	180
		cggccgtcaa ctatcgcaat		•	240
		cgagcatagt gtacgagaco			360
	ttaccctgcg	tgagggttgg gaatcagtca	a egetgetggg	Lygocotcae teccacegeg	360
			·		

gcgg	cgcc	tt a	catc	ggcg	c tc	cgct	tgaa	tcc	ctcc	gga	gtca	tgtg	ga t	ctga	tggta	
ggtg	ccgc	ta c	tgcg	tgct	c cg	ctct	ttac	atc	ggag	acc	tgtg	cggt	gg c	gtat	ttttg	I
gttg	gtca	ga t	gttc	tctt	t cc	agcc	gcgg	cgc	cact	gga	ctac	gcag	ga c	tgca	attgt	,
tcca	tcta	.cg c	9999	cacg	t ta	cggg	ccac	agg	atgg	ca						
<210 <211 <212 <213	> 1 > F	.88 .93 PRT lepat	itis	C V	irus											
<400	> 1	.88														
Thr 1	Cys	Gly	Phe	Ala 5	Asp	Leu	Met	Gly	Tyr 10	Ile	Pro	Leu	Val	Gly 15	Ala	
Pro	Val	Gly	Gly 20	Val	Ala	Arg	Ala	Leu 25	Ala	His	Gly	Val	Arg 30	Ala	Val	
Glu	Asp	Gly 35	Ile	Asn	Tyr	Ala	Thr 40	Gly	Asn	Leu	Pro	Gly 45	Cys	Ser	Phe	
Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys	Leu	Thr 60	Val	Pro	Thr	Ser	
Ala 65	Val	Asn	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Туг 75	His	Ile	Thr	Asn	Asp 80	
Cys	Pro	Asn	Ser	Ser 85	Ile	Val	Tyr	Glu	Thr 90	Glu	His	His	Ile	Leu 95	His	
Leu	Pro	Gly	Cys 100	Leu	Pro	Cys	Val	Arg 105	Val	Gly	Asn	Gln	Ser 110	Arg	Cys	
Trp	Val	Ala 115	Leu	Thr	Pro	Thr	Val 120	Ala	Ala	Pro	Tyr	Ile 125	Gly	Ala	Pro	
Leu	Glu 130		Leu	Arg	Ser	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr	
Ala 145		Ser	Ala	Leu	Tyr 150	Ile	Gly	Asp	Leu	Cys 155	Gly	Gly	Val	Phe	Leu 160	

Val Gly Gln Met Phe Ser Phe Gln Pro Arg Arg His Trp Thr Thr Gln 165 170 175

Asp Cys Asn Cys Ser Ile Tyr Ala Gly His Val Thr Gly His Arg Met 180 185 190

Ala

<210>	189	
<211>	579	
<212>	DNA	

<213> hepatitis C virus

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<210> 190

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 190

Thr Cys Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val 20 25 30

Glu Asp Gly Ile Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe

Ser	Ile 50	Phe	Leu	Leu	Ala	Leu 55	Leu	Ser	Cys		Thr 60	Val	Pro	Ala	Ser			
Ala 65	Glu	His	Tyr	Arg	Asn 70	Ala	Ser	Gly	Ile	Tyr 75	His	Ile	Thr	Asn	Asp 80			
Cys	Pro	Asn	Ser	Ser 85	Val	Val	Tyr	Glu	Thr 90	Asp	His	His	Ile	Leu 95	His			
Leu	Pro	Gly	Cys 100	Val	Pro	Cys	Val	Arg 105	Ala	Gly	Asn	Val	Ser 110	Arg	Сув			
Trp	Thr	Pro 115	Val	Thr	Pro	Thr	Val 120	Ala	Ala	Val	Ser	Met 125	Asp	Ala	Pro			
Leu	Glu 130	Ser	Phe	Arg	Arg	His 135	Val	Asp	Leu	Met	Val 140	Gly	Ala	Ala	Thr			
Val 145		Ser	Val	Leu	Tyr 150	Val	Gly	Asp	Leu	Cys 155	Gly	Gly	Ala	Phe	Leu 160			
Val	Gly	Gln	Met	Phe 165		Phe	Gln	Pro	Arg 170	Arg	His	Trp	Thr	Thr 175	Gln			
Asp	Cys	Asn	Cys 180		Ile	Tyr	Thr	Gly 185		Ile	Thr	Gly	His 190	Arg	Met		:	
Ala	L																	
<21 <21	.0> .1> .12> .13>	DNA	titi	s C	viru	ıs	·					•						
)0> agca		atco	taaa:	icc t	cáaa	gaaa	a ac	caaa	.cgta	aca	.ccaa	.ccg	ccgc	cccatg	60		
															cgcagg	1,20		
															.cgtggg	180		
															cccggg	240		
						aato										289		

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<210> 192 <211> 96 <212> PRT <213> hepatitis C virus	
<400> 192	
Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr As 1 5 10 15	sn
Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val G	ly
Gly Val Tyr Leu Leu Pro Arg Arg Gly Pro Arg Leu Gly Val Arg Al 35 40 45	la
Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60	ro
Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Pro G 65 70 75 86	ly 0
Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Tr 85 90 95	rp
<210> 193 <211> 498 <212> DNA <213> hepatitis C virus	
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ggcccccggt tgggtgtgcg cgcgactcgg aagacttcgg agcggtcgca acctcg	
aggegteaac etateceeaa ggegegeegg teegagggea ggteetggge geaage	cggg 240
tacccctggc ccctctatgg caatgagggc tgtgggtggg cagggtggct cctgtc	tcct 300
cgcggctctc ggccatcttg gggcccaaat gatccccggc ggagatcgcg caatct	gggt 360
aaggtcatcg ataccetgae gtgeggette geegaeetea tgggataeat eeeget	cgtg 420
ggcgcccccg tcgggggcgt cgccagggcc ctggcgcatg gcgtcagggc tgtgga	ggac 480

gggattaact atcgacag

498

<210> 194

<211> 166

<212> PRT

<213> hepatitis C virus

<400> 194

Met Ser Thr Asn Pro Lys Pro Gln Arg Lys Thr Lys Arg Asn Thr Asn 1 5 10 15

Arg Arg Pro Met Asp Val Lys Phe Pro Gly Gly Gly Gln Ile Val Gly 20 25 30

Gly Val Tyr Leu Leu Pro Arg Gly Pro Arg Leu Gly Val Arg Ala 35 40 45

Thr Arg Lys Thr Ser Glu Arg Ser Gln Pro Arg Gly Arg Arg Gln Pro 50 55 60

Ile Pro Lys Ala Arg Arg Ser Glu Gly Arg Ser Trp Ala Gln Ala Gly 65 70 75 80

Tyr Pro Trp Pro Leu Tyr Gly Asn Glu Gly Cys Gly Trp Ala Gly Trp 85 90 95

Leu Leu Ser Pro Arg Gly Ser Arg Pro Ser Trp Gly Pro Asn Asp Pro 100 105 110

Arg Arg Arg Ser Arg Asn Leu Gly Lys Val Ile Asp Thr Leu Thr Cys
115 120 125

Gly Phe Ala Asp Leu Met Gly Tyr Ile Pro Leu Val Gly Ala Pro Val 130 135 140

Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Ala Val Glu Asp 145 150 155 160

Gly Ile Asn Tyr Arg Gln 165

<210> 195

<211> 579

<212> DNA

<213> hepatitis C virus

<400> 195

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gggaatctgc	ctggttgctc	tttctctatc	ttcattcttg	cacttctctc	gtgcctcact	180
gtcccggcct	ctgcagttcc	ctaccgaaat	gcctctggga	tctatcatgt	caccaatgat	240
tgcccaaact	cttccatagt	ctatgaggca	gatgatctga	tcctacacgc	acctggctgc	300
gtgccttgtg	tcaggaaaga	taatgtgagt	aģgtgctggg	tccaaattac	ccccacgctg	360
tcagccccga	gcttcggagc	agtcacggct	ccccttcgga	gagccgttga	ttacttggtg	420
ggaggggctg	ccctctgctc	cgcgttatac	gttggagacg	cgtgtggggc	actatttttg	480
gtaggccaaa	tgttcaccta	taggcctcgc	cagcatgcta	cggtgcagga	ctgcaactgt	540
tccatctaca	gtggccacgt	caccggccat	cagatggca	•		579

<210> 196

<211> 193

<212> PRT

<213> hepatitis C virus

<400> 196

Thr Cys Gly Phe Ala Asp Leu Val Gly Tyr Ile Pro Leu Val Gly Gly
1 5 10 15

Pro Val Gly Gly Val Ala Arg Ala Leu Ala His Gly Val Arg Val Leu 20 25 30

Glu Asp Gly Val Asn Tyr Ala Thr Gly Asn Leu Pro Gly Cys Ser Phe 35 40 45

Ser Ile Phe Ile Leu Ala Leu Leu Ser Cys Leu Thr Val Pro Ala Ser 50 55 60

Ala Val Pro Tyr Arg Asn Ala Ser Gly Ile Tyr His Val Thr Asn Asp 65 70 75 80

Cys Pro Asn Ser Ser Ile Val Tyr Glu Ala Asp Asp Leu Ile Leu His 85 90 95

Ala Pro Gly Cys Val Pro Cys Val Arg Lys Asp Asn Val Ser Arg Cys
100 105 110

Trp Val Gln Ile Thr Pro Thr Leu Ser Ala Pro Ser Phe Gly Ala Val

115

Thr Ala Pro Leu Arg Arg Ala Val Asp Tyr Leu Val Gly Gly Ala Ala 135 Leu Cys Ser Ala Leu Tyr Val Gly Asp Ala Cys Gly Ala Leu Phe Leu 155 145 150 Val Gly Gln Met Phe Thr Tyr Arg Pro Arg Gln His Ala Thr Val Gln 170 165 Asp Cys Asn Cys Ser Ile Tyr Ser Gly His Val Thr Gly His Gln Met 185 180 Ala <210> 197 1485 <211> DNA <212> <213> hepatitis C virus <220> <221> misc feature <222> (787)..(787) "n" is any nucleotide <223> <220> <221> misc feature <222> (1011) . . (1011) <223> "n" is any nucleotide <220> <221> misc feature <222> (1444)..(1444) <223> "n" is any nucleotide <400> 197 tgtgccagga ccatcaccac cggagcttct atcacatact ccacttacgg caagttcctt 60 gctgatggag ggtgttcagg cggcgcgcat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga gctaggctcg tcgtcttggc cacggccacc cctcccggca gtgtgacaac gccccacccc 240 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt

ccccttgctt ttataaaggg tggtaggcat ctcatcttct gccattccaa gaaaaaatgt 360 gatgaactcg ccaagcaact gaccagcctg ggcgtgaacg ccgtggcata ttatagaggt 420 ctagacgtcg ccgtcatacc cacaacagga gacgtggtcg tgtgcagcac cgacgcgctc 480 atgacgggat tcaccggcga ctttgattct gtcatagact gcaactccgc cgtcactcag 540 acggtggact tcagtctgga tcccactttt accattgaga ctaccacagt gccccaggac 600 gcagtgtcca gaagccagcg ttggggccgc acggggagag gtaggcacgg catataccgg 660 tatgtctcgg ctggagagag accgtctggc atgttcgact ccgtggtgct ctgtgagtgc 720 tacgatgccg gatgtgcatg gtacgatctg actcctgccg agactaccgt gaggttgcgc 780 gcttacntaa acaccccgg gctccctgtc tgtcaggacc atttggaatt ctgggagggg 840 gtgttcacgg ggctcactaa catcgacgct cacatgctgt cacagaccaa acagggtggg 900 gagaatttcc cataccttgt agcgtaccaa gcaacagtct gtgttcgcgc gaaagcgccc 960 cccccagct gggacacaat gtggaaatgc atgctccgtc tcaaaccgac nttaactggc 1020 cctactcccc tcttgtacag gctggggccc gtccagaatg agatcacact gacgcacccc 1080 atcaccaagt acattatggc ttgcatgtct gcggacttgg aggtcattac cagcacttgg 1140 gttctggtgg ggggcgttgt ggcggccctg gcggcctact gcttgacggt gggttcggta 1200 gccatagtcg gtaggatcat cctctctggg aaacctgcca tcattcccga tagggaggta 1260 ttataccage aatttgatga gatggaggag tgctcggcct cgttgcccta tatggacgaa 1320 acacgtgcca ttgccggaca attcaaagag aaagtgctcg gcttcatcag cacgaccggc 1380 cagaaggctg aaactctgaa gccggcagcc acgtctgtgt ggaacaaggc tgagcagttc 1440 1485 tggnccacat acatgtggaa cttcatcagt gggatacaat aatag

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<210> 198
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<211> 484

<212> PRT

<213> hepatitis C virus

<220>

<221> MISC_FEATURE

<222> (263)...(263)

<223> "Xaa" is any amino acid

<220>

<221> MISC_FEATURE

<222> (337)..(337)

<223> "Xaa" is any amino acid

<220>

<221> MISC_FEATURE

<222> (482)..(482)

<223> "Xaa" is any amino acid

<400> 198

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Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala His Asp Val 20 25 30

Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly 35 40 45

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Thr Gly Asp Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160

Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175

Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190

- Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Trp 195 200 205
- Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220
- Gly Glu Arg Pro Ser Gly Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240
- Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255
- Val Arg Leu Arg Ala Tyr Xaa Asn Thr Pro Gly Leu Pro Val Cys Gln 260 270
- Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285
- Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Gly Glu Asn Phe Pro 290 295 300
- Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320
- Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335
- Xaa Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350
- Asn Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365
- Met Ser Ala Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly 370 375 380
- Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val 385 390 395 400
- Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415
- Asp Arg Glu Val Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser

425

420

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 440 435 Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 455 Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 475 470 Trp Xaa Thr Tyr <210> 199 <211> 1485 <212> DNA <213> hepatitis C virus <220> <221> misc_feature (684)..(684) <222> "n" is any nucleotide <223> <220> <221> misc feature <222> (909)..(909) <223> "n" is any nucleotide <220> <221> misc_feature <222> (1058)..(1058) <223> "n" is any nucleotide <220> <221> misc_feature (1133)..(1133)<222> "n" is any nucleotide <223> <400> 199 tgtgccagga ccatcaccac cggagettet atcacatact ccaettacgg caagtteett 60 gctgatggag ggtgttcagg cggcgcgtat gacgtgatca tatgcgacga gtgccattcc 120 caggacgcca ccaccattct tgggataggc actgtccttg accaggcaga gacggctgga 180 getaggeteg tegtettgge caeggecaec ceteceggea gtgtgaeaac geeceaecee 240 aacatcgagg aagtggccct gcctcaggag ggggaggttc ccttctacgg cagagccatt 300 ccccttgctt ttataaaggg tggtaggcat ctcatcttct gccattccaa gaaaaaatgt 360 420 gatgaactcg ccaagcaact gaccagcctg ggcgtgaacg ccgtggcata ttatagaggt ctagacgtcg ccgtcatccc cacagcagga gacgtggtcg tgtgcagcac cgacgcgctc 480 atgacgggat tcaccggcga ctttgattct gtcatagact gcaactccgc cgtcactcag 540 acggtggact tcagtctgga tcccactttt accattgaga ctaccacagt gccccaggac 600 gcagtgtcca gaagccagcg taggggccgc acggggagag gtaggcacgg catataccgg 660 tatgtctcgg ctggagagag accntctgac atgttcgact ccgtggtgct ctgtgagtgc 720 tacgatgccg gatgtgcgtg gtatgatctg actcctgccg agactaccgt gaggttgcgc 780 gcttacataa acaccccgg gctccctgtc tgtcaggacc atttggaatt ctgggagggg 840 gtgttcacgg ggctcactaa catcgacgct cacatgctgt cacagaccaa acagggtggg 900 gagaatttnc cataccttgt agcgtaccaa gcaacagtct gtgttcgcgc gaaagcgccc 960 cccccagct gggacacaat gtggaaatgc atgctccgtc tcaaaccgac tttaactggc 1020 cctactcccc tcttgtacag gctggggccc gtccagantg agatcacact gacgcacccc 1080 atcaccaagt acattatggc ttgcatgtct gcggacttgg aggtcattac cancacttgg 1140 gttctggtgg ggggcgttgt ggcggccctg gcggcctact gcttgacggt gggttcggta 1200 gccatagtcg gtaggatcat cctctctggg aaacctgcca tcattcccga tagggaggca 1260 ttataccagc aatttgatga gatggaggag tgctcggcct cgttgcccta tatggacgag 1320 acacgtgcca ttgccggaca attcaaagag aaagtgctcg gcttcatcag cacgaccggc 1380 cagaaggctg aaactctgaa gccggcagcc acgtctgtgt ggaacaaggc tgagcagttc 1440 1485 tgggccacat acatgtggaa cttcatcagc gggatacaat aatag

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<210> 200
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<211> 484

<212> PRT

<213> hepatitis C virus

<220>

<221> MISC FEATURE

<222> (228)..(228)

<223> "Xaa" is any amino acid

<220>

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<222> (303)..(303)

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<220>

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<223> "Xaa" is any amino acid

<220>

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<222> (378)..(378)

<223> "Xaa" is any amino acid

<400> 200

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Gly Lys Phe Leu Ala Asp Gly Gly Cys Ser Gly Gly Ala Tyr Asp Val 20 25 30

Ile Ile Cys Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly

Ile Gly Thr Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val 50 55 60

Val Leu Ala Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro 65 70 75 80

Asn Ile Glu Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr 85 90 95

Gly Arg Ala Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile 100 105 110

Phe Cys His Ser Lys Lys Lys Cys Asp Glu Leu Ala Lys Gln Leu Thr 115 120 125

Ser Leu Gly Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala 130 135 140

Val Ile Pro Thr Ala Gly Asp Val Val Val Cys Ser Thr Asp Ala Leu 145 150 155 160

- Met Thr Gly Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser 165 170 175
- Ala Val Thr Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile 180 185 190
- Glu Thr Thr Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg
 195 200 205
- Gly Arg Thr Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala 210 215 220
- Gly Glu Arg Xaa Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys 225 230 235 240
- Tyr Asp Ala Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr 245 250 255
- Val Arg Leu Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln 260 265 270
- Asp His Leu Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile 275 280 285
- Asp Ala His Met Leu Ser Gln Thr Lys Gln Gly Glu Asn Xaa Pro 290 295 300
- Tyr Leu Val Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro 305 310 315 320
- Pro Pro Ser Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro 325 330 335
- Thr Leu Thr Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln 340 345 350
- Xaa Glu Ile Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys 355 360 365
- Met Ser Ala Asp Leu Glu Val Ile Thr Xaa Thr Trp Val Leu Val Gly 370 380
- Gly Val Val Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val

390		395	400
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Ala Ile Val Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro 405 410 415

Asp Arg Glu Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser 420 425 430

Ala Ser Leu Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe 435 440 445

Lys Glu Lys Val Leu Gly Phe Ile Ser Thr Thr Gly Gln Lys Ala Glu 450 455 460

Thr Leu Lys Pro Ala Ala Thr Ser Val Trp Asn Lys Ala Glu Gln Phe 465 470 475 480

Trp Ala Thr Tyr

<210> 201

385

<211> 340

<212> DNA

<213> hepatitis C virus

<400> 201

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tatgtacaat agcaagggag acctttgcgg gtatcgcagg tgccgcgcaa gcggcgtata 180
taccaccagc ttcgggaaca cactgacgtg ctaccttaaa gcctcagcag ccatcagggc 240
tgcggggctg aaggactgca ccatgctggt ttgcggtgac gacttagtcg tgatcgctga 300
aagcggtggc gtcgaggagg acaagcgagc cctcggagct 340

<210> 202

<211> 113

<212> PRT

<213> hepatitis C virus

<400> 202

Ser Thr Val Thr Glu Arg Asp Ile Arg Val Glu Glu Glu Val Tyr Gln 1 5 10 15

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Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Tyr	Asn	Ser	Lys 45	Gly	Asp	Leu	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe	
Gly 65	Asn	Thr	Leu	Thr	Cys 70	Tyr	Leu	Lys	Ala	Ser 75	Ala	Ala	Ile	Arg	Ala 80	
Ala	Gly	Leu	Lys	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95		
Val	Ile	Ala	Glu 100	Ser	Gly	Gly	Val	Glu 105	Glu	Asp	Lys	Arg	Ala 110	Leu	Gly	٠
Ala				-	•											
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Cys	Cys	Asp	Leu 20	Glu	Pro	Glu	Thr	Arg 25	Lys	Val	Ile	Ser	Ala 30	Leu	Thr	
Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	His	Asn	Ser	Arg 45	Gly	Asp	Leu	
Cys	Gly 50	Tyr	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe	
Gly 65	Asn	Thr	Leu	Thr	Cys 70	Phe	Leu	Lys	Ala	Thr 75	Ala	Ala	Thr	Lys	Ala 80	
Ala	Gly	Leu	Lys	Asp 85	Cys.	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
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cgc	aggt	.ctg	aaag	Igttg	rca g	catg	ctgg	rt tt	gtgg	agac	gac	ttag	tcg	tcat	ctgcga	300
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Glu	Arg	Leu 35	Tyr	Val	Gly	Gly	Pro 40	Met	Tyr	Asn	Ser	Lys 45	Gly	Asp	Leu		٠
Cys	Gly 50	Gln	Arg	Arg	Cys	Arg 55	Ala	Ser	Gly	Val	Tyr 60	Thr	Thr	Ser	Phe		
Gly 65	Asn	Thr	Val	Thr	Cys 70	Tyr	Leu	Lys	Ala	Val 75	Ala	Ala	Thr	Arg	Ala 80		
Ala	Gly	Leu	Lys	Gly 85	Cys	Ser	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val		
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Glu Arg Leu Tyr Lys Gly Gly Pro Met Tyr Asn Ser Lys Gly Asp Leu 35 40 45

Cys Gly Leu Arg Arg Cys Arg Ala Ser Gly Val Tyr Thr Thr Ser Phe 50 55 60

Gly Asn Thr Val Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 65 70 75 80

Ala Gly Leu Lys Asp Cys Thr Met Leu Val Cys Gly Asp Asp Leu Val 85 90 95

Val Ile Ala Glu Ser Gly Gly Val Glu Glu Asp Ala Arg Ala Leu Arg 100 105 110

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cgcgggcttg aaggattgca ccatgctggt ctgcggngac gacctggttg tcattgctga
                                                                     300
                                                                     340
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Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala 75 70

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Val Ile Ala Glu Ser Ile Gly Ile Asp Glu Asp Lys Gln Ala Leu Arg 100 105

Thr

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Cys Gly Gln Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Phe
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Gly Asn Thr Leu Thr Cys Tyr Leu Lys Ala Thr Ala Ala Thr Arg Ala

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120

180

240

300

340

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Pro

213		
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DNA		`
hepatitis	C	virus
	340 DNA *	340

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Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Leu Thr Thr Ser Cys

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Ala	Lys	Leu	Gln	Asp 85	Cys	Thr	Met	Leu	Val 90	Cys	Gly	Asp	Asp	Leu 95	Val	
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gag	ccaç	gggt	aacg	agga	igg a	cgag	gcgaa	a co	tgag	gagct	:					340
	.1> l2> l3>	216 113 PRT hepa	atiti	is C	vir	ıs										
		r Val	l Thi	r Gli 5	u Ar	g As	p Ile	e Ar	g Thi	r Gli	u Glu	ı Sei	r Ile	e Ty:	r Gln	
Ala	а Су	s Se:	r Let 20	u Pr	o Gl	n Gl	u Ala	a Ar 25	g Th	r Vai	1 11	e His	30	r Le	u Thr	
Gl	u Ar	g Le [.] 35	u Ty:	r Va	1 G1	y Gl	y Pr 40		t Il	e As:	n Se	r Ly: 45	s Gl	y Gl	n Ser	

Cys Gly Tyr Arg Arg Cys Arg Ala Ser Gly Val Phe Thr Thr Ser Met 50 55 60

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gag	tgat	ggc	gtcg	acga	.gg a	taga	.gcaç	ic cc	tgag	agcc	!			1		340
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Суя	в Суя	s Ası	n Lei 20	ı Glı	ı Pro	o Gli	ı Ala	a Arg 25	g Ly:	s Val	l Ile	e Sei	30	r Lei	ı Thr	
Gl.	u Ar	g Let 35	u Tyi	r Cy	s Gl	y Gl	y Pr 40	o Me	t Ph	e Ası	n Se	r Ly: 45	s Gl	y Ala	a Gln	
Су	s Gl	у Ту:	r Ar	g Ar	д Су	s Arg	g Al	a Se	r Gl	y Va	l Le 60	u Pro	o Th:	r Se	r Phe	

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Trp Asp Glu Met Trp Lys Cys Leu Val Arg Leu Lys Pro Thr Leu His
50 55 60

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 65 70 75 80

Cys Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 85 90 95

Asp Leu Glu Val Thr Thr Ser Thr Trp Val Leu Leu Gly Gly Val Leu 100 105 110

Ala Ala Leu Ala Ala Tyr Cys Leu Ser Val Gly Cys Val Val Ile Val 115 120 125

Gly His Ile Glu Leu Gly Gly Lys Pro Ala Ile Val Pro Asp Lys Glu 130 135 140

Val Leu Tyr Gln Gln Tyr Asp Glu Met Glu Glu Cys Ser Gln Ala Ala 145 150 155 160

Pro Tyr Ile Glu Gln Ala Gln Val Ile Ala His Gln Phe Lys Glu Lys 165 170 175

Val Leu Gly Leu Leu Gln Arg Ala Thr Gln Gln Ala Val Ile Glu 180 185 190

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Asp Glu Cys His Ser Gln Asp Ala Thr Thr Ile Leu Gly Ile Gly Thr 35 40 45

Val Leu Asp Gln Ala Glu Thr Ala Gly Ala Arg Leu Val Val Leu Ala
50 55 60

Thr Ala Thr Pro Pro Gly Ser Val Thr Thr Pro His Pro Asn Ile Glu 65 70 75 80

Glu Val Ala Leu Pro Gln Glu Gly Glu Val Pro Phe Tyr Gly Arg Ala 85 90 95

Ile Pro Leu Ala Phe Ile Lys Gly Gly Arg His Leu Ile Phe Cys His
100 105 110

Ser Lys Lys Lys Cys Asp Glu Léu Ala Lys Gln Leu Thr Ser Leu Gly 115 120 125

Val Asn Ala Val Ala Tyr Tyr Arg Gly Leu Asp Val Ala Val Ile Pro 130 135 140 Thr Ala Gly Asp Val Val Cys Ser Thr Asp Ala Leu Met Thr Gly
145 150 155 160

Phe Thr Gly Asp Phe Asp Ser Val Ile Asp Cys Asn Ser Ala Val Thr 165 170 175

Gln Thr Val Asp Phe Ser Leu Asp Pro Thr Phe Thr Ile Glu Thr Thr 180 185 190

Thr Val Pro Gln Asp Ala Val Ser Arg Ser Gln Arg Arg Gly Arg Thr 195 200 205

Gly Arg Gly Arg His Gly Ile Tyr Arg Tyr Val Ser Ala Gly Glu Arg 210 215 220

Pro Ser Asp Met Phe Asp Ser Val Val Leu Cys Glu Cys Tyr Asp Ala 225 230 235 240

Gly Cys Ala Trp Tyr Asp Leu Thr Pro Ala Glu Thr Thr Val Arg Leu 245 250 255

Arg Ala Tyr Ile Asn Thr Pro Gly Leu Pro Val Cys Gln Asp His Leu 260 265 270

Glu Phe Trp Glu Gly Val Phe Thr Gly Leu Thr Asn Ile Asp Ala His 275 280 285

Met Leu Ser Gln Thr Lys Gln Gly Gly Glu Asn Phe Pro Tyr Leu Val 290 295 300

Ala Tyr Gln Ala Thr Val Cys Val Arg Ala Lys Ala Pro Pro Pro Ser 305 310 315 320

Trp Asp Thr Met Trp Lys Cys Met Leu Arg Leu Lys Pro Thr Leu Thr 325 330 335

Gly Pro Thr Pro Leu Leu Tyr Arg Leu Gly Pro Val Gln Asn Glu Ile 340 345 350

Thr Leu Thr His Pro Ile Thr Lys Tyr Ile Met Ala Cys Met Ser Ala 355 360 365

Asp Leu Glu Val Ile Thr Ser Thr Trp Val Leu Val Gly Gly Val Val 370 380

Ala Ala Leu Ala Ala Tyr Cys Leu Thr Val Gly Ser Val Ala Ile Val 385 390 395 400

Gly Arg Ile Ile Leu Ser Gly Lys Pro Ala Ile Ile Pro Asp Arg Glu 405 410 415

Ala Leu Tyr Gln Gln Phe Asp Glu Met Glu Glu Cys Ser Ala Ser Leu 420 425 430

Pro Tyr Met Asp Glu Thr Arg Ala Ile Ala Gly Gln Phe Lys Glu Lys 435 440 445

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Ser Lys Gln Ala 20

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Leu Lys Ser Lys 20

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